

OIPE

RAW SEQUENCE LISTING DATE: 05/04/2000
 PATENT APPLICATION: US/09/549,848 TIME: 15:30:39

Input Set : A:\Seqlist.txt
 Output Set: N:\CRF3\05042000\I549848.raw

4 <110> APPLICANT: Lassner, M
 5 Post-Beittenmiller, D
 6 Savidge, B
 7 Weiss, J
 9 <120> TITLE OF INVENTION: Nucleic Acid Sequences Involved in
 10 Tocopherol Synthesis
 12 <130> FILE REFERENCE: 17133/02
 14 <140> CURRENT APPLICATION NUMBER: US/09/549,848
 14 <141> CURRENT FILING DATE: 2000-04-14
 14 <150> PRIOR APPLICATION NUMBER: 60/129,899
 15 <151> PRIOR FILING DATE: 1999-04-15
 17 <150> PRIOR APPLICATION NUMBER: 60/146,461
 18 <151> PRIOR FILING DATE: 1999-07-30
 20 <160> NUMBER OF SEQ ID NOS: 94
 22 <170> SOFTWARE: FastSEQ for Windows Version 4.0

ERRORED SEQUENCES

1599 <210> SEQ ID NO: 90
 E--> 1600 <211> LENGTH 0
 1601 <213> ORGANISM: Artificial Sequence

1603 <400> SEQUENCE: 90
 1604 gaattcactt caaaaaaggt aacag

(mandatory response)

see item 12 on Error Summary Sheet

**Does Not Comply
 Corrected Diskette Needed**

*25 ← insert
 cumulative base total at right margin
 of each line*

✓ FYI

Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

VERIFICATION SUMMARY

DATE: 05/04/2000

PATENT APPLICATION: US/09/549,848

TIME: 15:30:40

Input Set : A:\Seqlist.txt

Output Set: N:\CRF3\05042000\I549848.raw

L:14 M:270 C: Current Application Number differs, Replaced Current Application No
L:14 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:311 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:312 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:676 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:677 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:678 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:757 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:770 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:787 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:788 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:790 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27
L:1600 M:210 E: (40) Invalid Number of Sequences, LENGTH:
L:1603 M:282 W: Numeric Field Identifier Missing, <212> is required.